

## AMENDMENTS TO THE SPECIFICATION

**Please replace Paragraph [0015] of the specification with the following paragraph:**

[0015] Figure 5 shows human (SEQ ID NO: 1), rat (SEQ ID NO: 2), and mouse (SEQ ID NO: 3) sequences for Dipeptidyl Peptidase IV during intestinal differentiation which is also useful in assays of preferred embodiments.

**Please replace Paragraph [0068] of the specification with the following paragraph:**

[0068] The presence of anti-Purkinje cell antibodies in some PCD patients suggests an autoimmune etiology. To identify the molecular targets for these autoantibodies, an Agt1 1 cDNA expression library from human cerebellum was constructed and screened with IgG from a patient with paraneoplastic cerebellar degeneration. A single clone, pCDR2, produced a fusion protein that reacted strongly with the patient's IgG. Sequencing the pCDR clones revealed 6 amino acids repeated in tandem along the entire cDNA sequence (VAL, PRO, LEU, LEU, GLU, ASP) (SEQ ID NO:4). This gene was expressed predominantly in neuroectodermal tissues (68).

**Please replace Paragraph [0078] of the specification with the following paragraph:**

[0078] The following antigens, proteins, peptides, enzymes, tissue receptors, lymphocyte receptors, neurotransmitters listed below are representative of antigens used in assays of preferred embodiments.

MBP Sequence 87-106	VVHFFKNIVTPRTPPPSQGK <u>(SEQ ID NO:5)</u>
MBP Sequence 83-89	ENPVVHFFKNIVTPRTP <u>(SEQ ID NO:6)</u>
MBP Sequence 1-11	ASQKRPSQRSK <u>(SEQ ID NO:7)</u>
MBP Sequence 200-211	ANMQRQAVPTL <u>(SEQ ID NO:8)</u>
Proteolipid Protein Sequence 40-60	TGTEKLIETYFSKNYQDYEYL <u>(SEQ ID NO:9)</u>
Proteolipid Protein Sequence 89-106	GFYTTGAVRQIFGDYKTT <u>(SEQ ID NO:10)</u>
Proteolipid Protein Sequence 103-120	YKTTICGKGLSATVTGGQ <u>(SEQ ID NO:11)</u>
Proteolipid Protein Sequence 125-143	SRGQHQAHSLERVCHCLGK <u>(SEQ ID NO:12)</u>
Proteolipid Protein Sequence 139-154	HCLGKWLGHDPDKFVGI <u>(SEQ ID NO:13)</u>
Transaldolase Protein Sequence 11-25	MESALDQLKQFTTVV <u>(SEQ ID NO:14)</u>
Transaldolase Protein Sequence 21-35	ETTVVADTGDFHAID <u>(SEQ ID NO:15)</u>

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Transaldolase Protein Sequence 31-45      FHAIDEYKPQDATTN (SEQ ID NO:16)  
Transaldolase Protein Sequence 71-85      KLGGSEQEDQIKNAID (SEQ ID NO:17)  
Transaldolase Protein Sequence 81-95      KNAIDKLFVLFGAEI (SEQ ID NO:18)  
Transaldolase Protein Sequence 261-275      GELLQDNAKLVPVLS (SEQ ID NO:19)  
Transaldolase Protein Sequence 271-285      VPVLSAKAAQASDLE (SEQ ID NO:20)  
Transaldolase Protein Sequence 311-325      GIRKFAADAVKLERM (SEQ ID NO:21)  
MOG Sequence 1-20      GQFRVIGPRHPIRALVGDEV (SEQ ID NO:22)  
MOG Sequence 61-80      QAPEYRGRTELLKDAIGEGK (SEQ ID NO:23)  
MOG Sequence 101-120      RDHSYQEEAAMELKVEDPFY (SEQ ID NO:24)  
MOG Sequence 145-160      VFLCLQYRLRGKLRAE (SEQ ID NO:25)  
MAG Sequence 37-60REIVDRKYSICKSGCFYQKKEEDW (SEQ ID NO:26)  
Sodium Ion Channel Na 1.2      TVTVPIALGESDFENLNTEEFSSSESDM (SEQ ID NO:27)  
                                 Na 1.3      TVTVPIAVGESDFENLNTEEFSSSESEL (SEQ ID NO:28)  
                                 Na 1.1      TVTVPIAVGESDFENLNTEDFSSSESDL (SEQ ID NO:29)  
                                 Na 1.6      TVRVPIAVGESDFENLNTEDVSSESDP  
                                 (SEQ ID NO:30)  
Glutamate Receptor      ANEYERFVPFSDQQISNDAAC (SEQ ID NO:31)  
Cerebellar peptides      FLEDVPLLEDIPLLEDVPLLED (SEQ ID NO:32)  
                                 FLEDVPLLEDIPLLEDVP (SEQ ID NO:33)  
                                 LLEDTDFLEDPDFLEAID (SEQ ID NO:34)  
Amyloid  $\beta$       DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGGVIA  
                                 (SEQ ID NO:35)  
CD69-Human      MECEKNLYWICNKPYK (SEQ ID NO:36)  
Zinc Finger Protein      PYKCPECCKSFSQKSDLVKHQRTHTG (SEQ ID NO:37)  
Glucose Regulated Protein-78 (GRP-78)      EEEDKKEDVGTVVGI (SEQ ID NO:38)  
Vasoactive Intestinal Peptide      NYTRLRKQMAVKKYL (SEQ ID NO:39)  
Gliadin Peptides      QPFRPQQPYQPQPQYSQPQQ (SEQ ID NO:40)  
                                 QPYQPQPQYSQPQQPISQQQ (SEQ ID NO:41)  
                                 QFLGQQQPFPPQPYQPQPQPF (SEQ ID NO:42)  
                                 PLVQQQQFLGQQQPFPPQPY (SEQ ID NO:43)

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HNVVHAILHQQQQQQQEQKQ (SEQ ID NO:44)  
NPSQQQPQEQVPLVQQQ (SEQ ID NO:45)  
QQLPQPQQPQQSFPQQQPF (SEQ ID NO:46)  
Gluteomorphin YPFPGPIP (SEQ ID NO:47)  
Casomorphin GYYPTYGGWL (SEQ ID NO:48)  
Secretin (human) HSDGTFTSELSRLREGARLQRLQLV (SEQ ID NO:49)  
Campylobacter Jejuni Toxin TPPLLAAILMLASLRSHIVSDHFPVNFRKF (SEQ ID NO:50)  
 $\alpha$ -S1 Casein  
RPKHPIKHQGLPQEVLENENLLRFFVAPFPEVFGKEKVNELSKDIGSESTDEQAMED  
IKQMEAESISSSEEIVPNSVEQKHIQKEDVPSELYLGYLEQLRLKKYKVPQLEIVP  
NSAEERLHSMKEGIHAQQKEPMIGVNQELAYFYPELFRQFYQLDAYPSGAWYYV  
PLGTQYTDAPSFSDIPNPIGSENSEKTTMPLW (SEQ ID NO:51)  
 $\alpha$ -S2 Casein MKEGIHAQQK (SEQ ID NO:52)  
YQKFALPQYL (SEQ ID NO:53)  
K Casein KDERFFSDKI (SEQ ID NO:54)  
SPPEINTVQV (SEQ ID NO:55)  
Vasoactive Intestinal Peptide HSDAVFTDNYTRLRKQMAVKKYLNSILN  
(SEQ ID NO:56)  
Somatostatin YSANSNPAMAPRERKAGCKNFFWKFTTSC (SEQ ID NO:57)  
Substance P RQKPQQFFGLM (SEQ ID NO:58)  
Oxytocin CYKQNCPLG (SEQ ID NO:59)  
Pancreatic Peptide APLEPVYPGDNATPEQMAQYAADLRRYINMLTRPRY  
(SEQ ID NO:60)  
Gastrin-1 EGPWLEEEEEAYGWMD (SEQ ID NO:61)  
Big Gastrin-1 ELGPQGPPHLVADPSKKQGPWLEEEEEAYGWMD (SEQ ID NO:62)  
Gastrin Releasing Peptide VPLPAGGGTVLTKMYPRGNHWAVGHLM (SEQ ID NO:63)  
Enkephalin YGGFLM (SEQ ID NO:64)  
 $\beta$ -Endorphin YGGFMTSEKSQTPLVTLFKNAIKNAYKKGE (SEQ ID NO:65)  
Big Endorphin CSCSSLMDKECVYFCHLDIIWVNTPEHVVPYGLGSPRS  
(SEQ ID NO:66)

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Dynorphin A YGGFLRRIRPKLKWDNQ (SEQ ID NO:67)  
Dynorphin B YGGFLRRQFKVVT (SEQ ID NO:68)  
Serotonin Receptor MPHLLSGFLEV TASPAPTWDAP (SEQ ID NO:69)  
IFGHFFCNVFIAMDVMCCTASI (SEQ ID NO:70)  
LKLAERPERSEFVLQNSDHCGK (SEQ ID NO:71)  
Fibrillin SFRPGSRGGSRG (SEQ ID NO:72)  
Calreticullin EQFLDGDGWTSRWIESGLQTSQ (SEQ ID NO:73)  
Motillin FVPIFTY GELQRMQEKERNKGQ (SEQ ID NO:74)  
Chlamydia HSP-60 LKQIAAHAGKEGAIIFQQVM (SEQ ID NO:75)  
Human HSP-60 1-20 MLRLPTVFRQMRPVSRVLAP (SEQ ID NO:76)  
16-35 RVLAPHLTRAYAKDVKFGAD (SEQ ID NO:77)  
31-50 KFGADARALMLQGVDLLADA (SEQ ID NO:78)  
46-65 LLADAVAVTMGPKGRTVIE (SEQ ID NO:79)  
61-80 TVIEEQSWGSPKVT KDGVTV (SEQ ID NO:80)  
76-95 DGVTVAKSIDLKDKYKNIGA (SEQ ID NO:81)  
91-110 KNIGAKLVQDVANNTNEEAG (SEQ ID NO:82)  
106-125 NEEAGDGT TATV LARSIK (SEQ ID NO:83)  
121-140 RSIKKEGFEKISKGANPVEI (SEQ ID NO:84)  
136-155 NPVEIRRGVMLAVDAVIAEL (SEQ ID NO:85)  
151-170 VIAELKKQSKPVTTP EEIAQ (SEQ ID NO:86)  
166-185 EEIAQVATISANGDKEIGNI (SEQ ID NO:87)  
181-199 EIGNISDAMKKVGRKGVI (SEQ ID NO:88)  
195-214 RKGVITVKDGKTLNDELEII (SEQ ID NO:89)  
210-229 ELEII EGMKFDRGYISPYFI (SEQ ID NO:90)  
225-244 SPYFINTSKGQKCEFQDAYV (SEQ ID NO:91)  
240-259 QDAYVLLSEKKISSIQSIVP (SEQ ID NO:92)  
255-275 QSIVPALEIANAH RKPLVIA (SEQ ID NO:93)  
271-290 LVIIAEDVDGEALSTLV LNR (SEQ ID NO:94)  
286-305 LVLNRLKVGLQVVAVKAPGF (SEQ ID NO:95)  
301-320 KAPGFGDNRKNQLKDMAIAT (SEQ ID NO:96)

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316-335 MAIATGGAVFGEEGLTLNLE (SEQ ID NO:97)  
331-350 TLNLEDVQPHDLGKVGEVIV (SEQ ID NO:98)  
346-365 GEVIVTKDDAMLLKGKGDKA (SEQ ID NO:99)  
361-380 KGDKAQIEKRIQEIEQLDV (SEQ ID NO:100)  
376-395 EQLDVTTSEYEKEKLNERLA (SEQ ID NO:101)  
391-410 NERLAKLSDGVAVLKVGGTS (SEQ ID NO:102)  
406-425 VGGTDVEVNEKKDRVTDAL (SEQ ID NO:103)  
421-440 VTDALNATRAAVEEGIVLGG (SEQ ID NO:104)  
436-455 IVLGGGCALLRCIPALDSLT (SEQ ID NO:105)  
451-470 LDSLTPANEDQKIGIEIKR (SEQ ID NO:106)  
466-485 EIIKRTLKIPAMTIKNAGV (SEQ ID NO:107)  
481-500 KNAGVEGSLIVEKIMQSSSE (SEQ ID NO:108)  
496-515 QSSSEVGYDAMAGDFVNMVE (SEQ ID NO:109)  
511-530 VNMVEKGIIDPTKVVRTALL (SEQ ID NO:110)  
526-545 RTALLDAAGVASLLTTAEVV (SEQ ID NO:111)  
541-560 TAEVVVTEIPKEEKDPGMGA (SEQ ID NO:112)  
556-573 PGMGAMGGMGGGMGGGMF (SEQ ID NO:113)  
437-460 VLGGGVLLLRVIPALDSLTPANED (SEQ ID NO:114)

Dipeptidylpeptidase peptides

Peptide 1 MKTPWRVLLGLLGAAALVTIITVPVLLNK  
(SEQ ID NO:115)  
Peptide 2 MAEYGNSSVFLENSTFDEFGH (SEQ ID NO:116)  
Peptide 3 KRQLITEERIPNNTQWVTWSP (SEQ ID NO:117)  
Peptide 4 NGTFLAYAQFNDTEVPLIEYS (SEQ ID NO:118)  
Peptide 5 VTNATSIQITAPASMLIGDHY (SEQ ID NO:119)  
Peptide 6 IQNYSVMDICDYDESSGRWNC (SEQ ID NO:120)  
Peptide 7 NSFYKIISNEEGYRHICYFQI (SEQ ID NO:121)  
Peptide 8 NVQMPSKKLDFILNETKFWY (SEQ ID NO:122)  
Peptide 9 PEDNLDHYRNSTVMSRAENFK (SEQ ID NO:123)  
Peptide 10 TAHQHIYTHMSHFQKQCFSLP (SEQ ID NO:124)

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Figure 5 shows sequences for Dipeptidyl Peptidase IV during intestinal differentiation which is also useful in assays of preferred embodiments. Figure 5 compares the amino acid sequences of human, rat, and mouse DPP IV, respectively. These sequences are aligned. The potential sites for phosphorylation (T or S) and for N-glycosylation (NXT) are displayed as underlined. Preferred embodiments include the sequences listed above, along with counterparts which have post-translational modifications.

**Please replace Paragraph [0122] of the specification with the following paragraph:**

[0122] Gliadin peptides QQLPQPQQPQQSFPQQQPF (SEQ ID NO:125), LQLQFPQPQLPYPQPQLPY (SEQ ID NO:126) - P Q P L P Y P Q P Q P F (SEQ ID NO:127), QQPQQFZPQQPYYPZXPZLGZZZPFPPZ (SEQ ID NO:128), gluteomorphin ZGZPGYYPTSPZZPGQEQ (SEQ ID NO:129), casomorphin ZTZSLVYFPFGPIPNSLP (SEQ ID NO:130), B-casein LHLPLLLZSWMHZPHZPL (SEQ ID NO:131) and CD69 antibody binding epitope MECEKNLYWICNKPYK (SEQ ID NO:132) were synthesized by Bio-Synthesis Inc. (Lewisville, TX). Dipeptidylpeptidase IV (CD26), streptokinase (SK), lipopolysaccharide (LPS), human serum albumin (HSA), mercury [o-carboxyphenyl] Thio] ethyl mercury sodium salt (Thimerosal) were purchased from Sigma (St. Louis, MO).

**Please replace Paragraph [0136] of the specification with the following paragraph:**

[0136] *Gliadin peptides:* Gliadin peptide QQLPQPQQPQQSFPQQQPF (SEQ ID NO:125) and *Chlamydia trachomatis* HSP-60 peptide LKQIAAHAGKEGAIIFQQVM (SEQ ID NO:133), HPLC grade, were synthesized by Bio-Synthesis Inc. (Lewisville, TX).

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### **AMENDMENTS TO THE DRAWINGS**

Enclosed please find corrected Figure 5 with red ink markings designating the proposed changes to the drawing in this application for which approval by the Examiner is requested. Also enclosed is an amended version of Figure 5. Figure 5 has been amended to add sequence identification numbers 1-3 to the sequence.